Cellular Expression of β_2 AR- β gal $\Delta\alpha$ Fusion Protein in C2 Clones (measured by anti- β -gal ELISA)

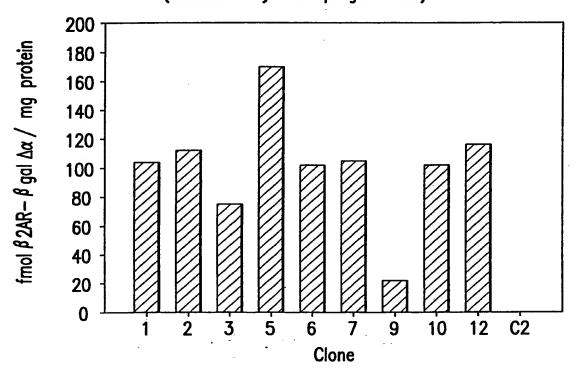


FIG. 1A



Cellular expression of β Arr- β gal $\Delta\omega$ fusion protein in C2 clones (measured by anti- β gal ELISA)

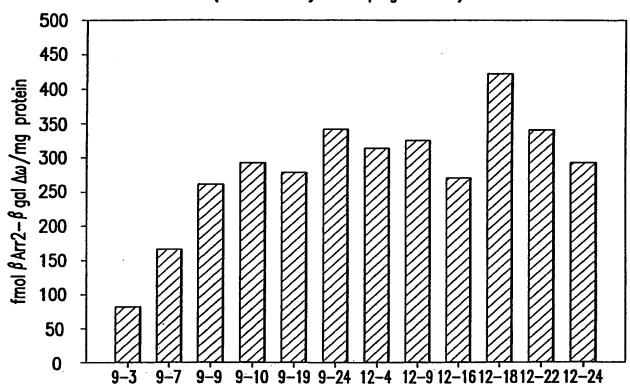


FIG. 1B



Agonist Stimulated cAMP Response in C2 Cells Expressing $\beta 2AR - \beta gal \Delta \alpha$

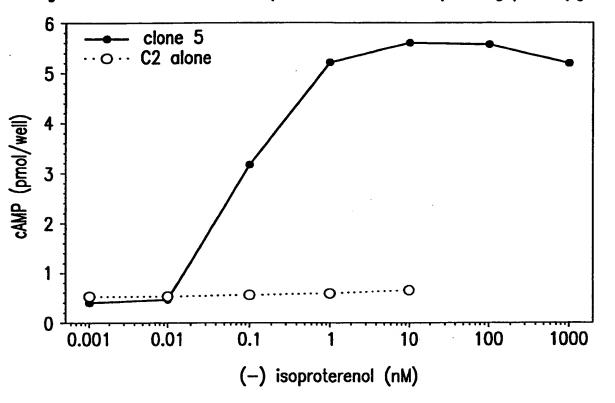


FIG.2

β -galactosidase Complementation as a Measurement for β_2 AR- β gal $\Delta\alpha$ interacting with β Arrestin2- β gal $\Delta\omega$ upon agonist Stimulation

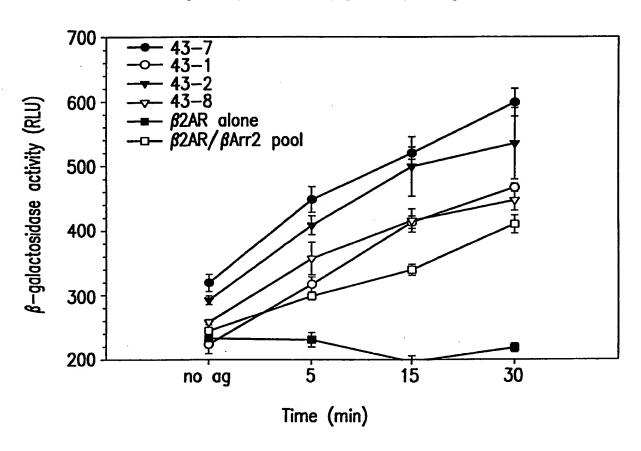
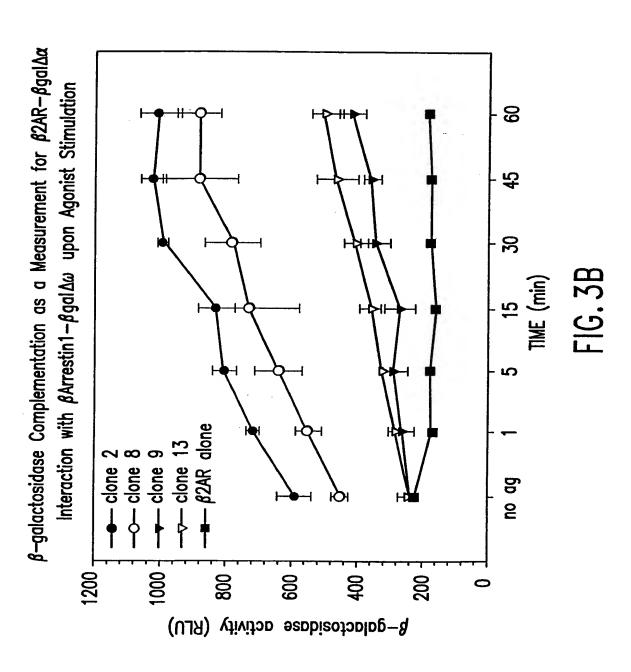


FIG. 3A







β -galactosidase Activity in Response to Agonist in C2 Cells Coexpressing β 2AR- β gal $\Delta\alpha$ and β Arrestin2- β gal $\Delta\omega$ Fusion Proteins

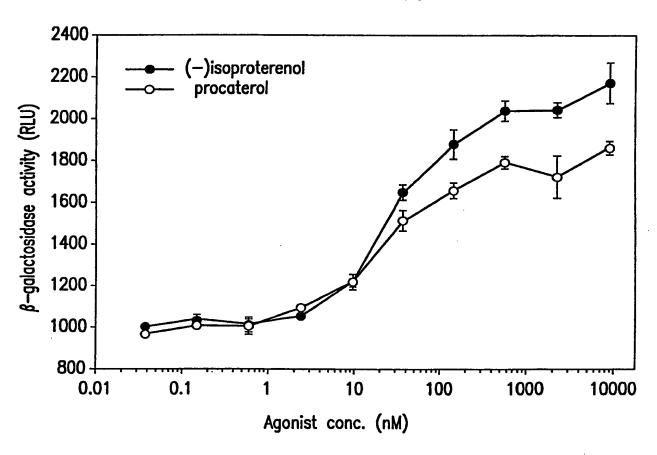


FIG. 4A



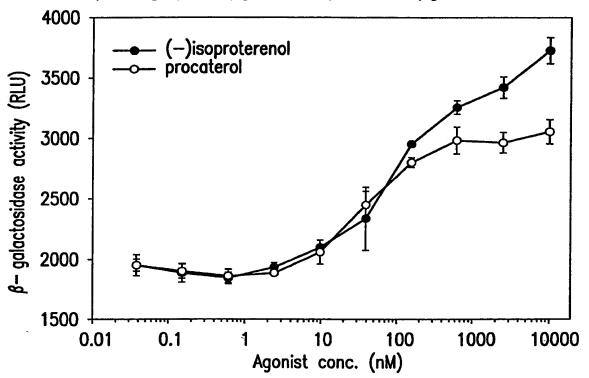


FIG. 4B

Inhibition of β -galactosidase activity in C2 Cells Coexpressing β 2AR $-\beta$ gal $\Delta\alpha$ and β Arrestin2- β gal $\Delta\omega$ Fusion Proteins

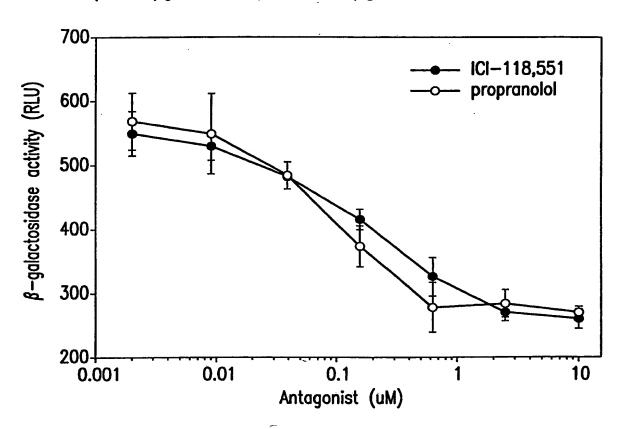


FIG. 5A

Antagonist Inhibition of β -galactosidase Activity in C2 Cells Coexpressing $\beta 2AR-\beta gal\Delta\alpha$ and $\beta Arrestin 1-\beta gal\Delta\omega$ Fusion Proteins

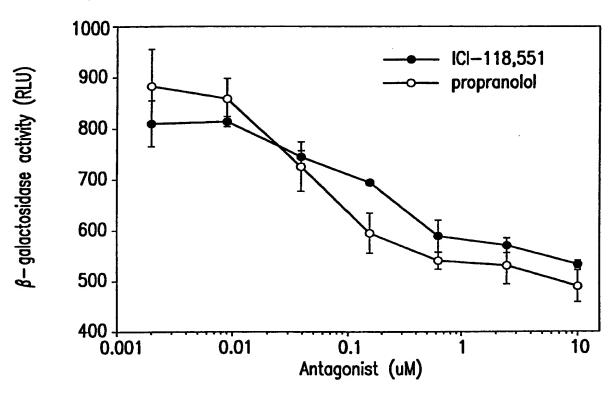


FIG. 5B



Agonist Stimulated cAMP Response in Clones or Pools of C2 Cells Coexpressing A2aR- β gal $\Delta\alpha$ and β Arrestin1- β gal $\Delta\omega$ Fusion Proteins

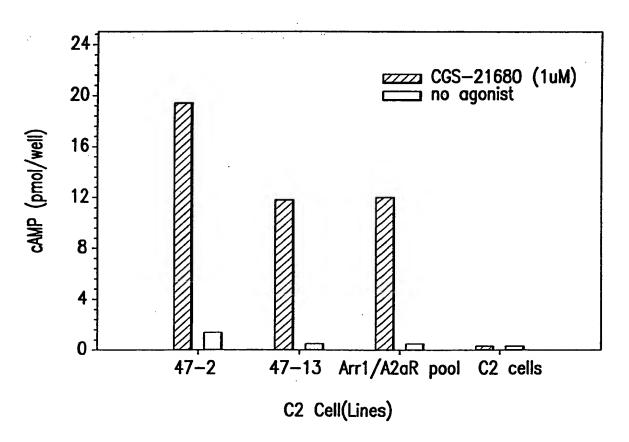


FIG.6

Agonist Stimulated cAMP Response in Clones or Pools of C2 Cells Expressing D1- β gal $\Delta\alpha$ and β Arrestin2- β gal $\Delta\omega$ Fusion Proteins

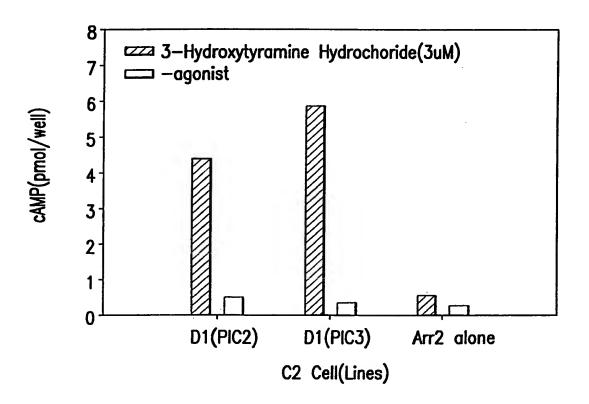


FIG. 7



 β_2 AR- β gal $\Delta\omega$ and β arr2- β gal $\Delta\alpha$ Interaction in HEK293 Clones in Response to Isoproterenol Treatment (1 μ M)

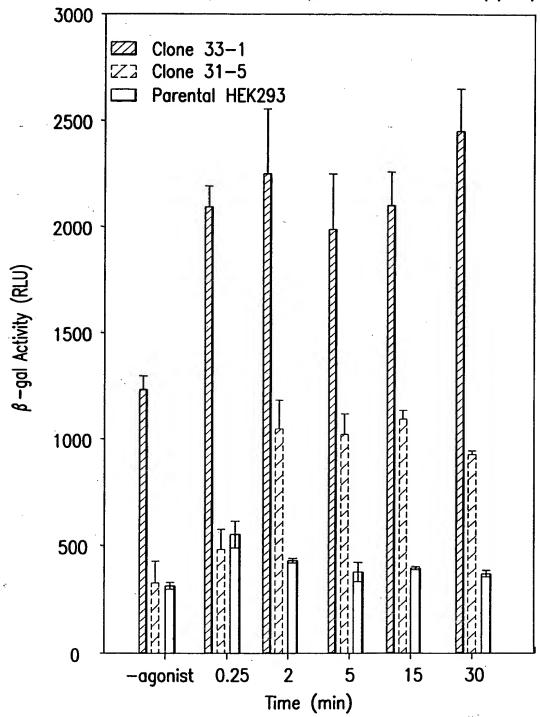


FIG. 8A

 β 2AR- β gal $\Delta\alpha$ and β Arr1- β gal $\Delta\omega$ Interaction in a CHO Pool in Response to Isoproterenol Treatment(10 μ M)

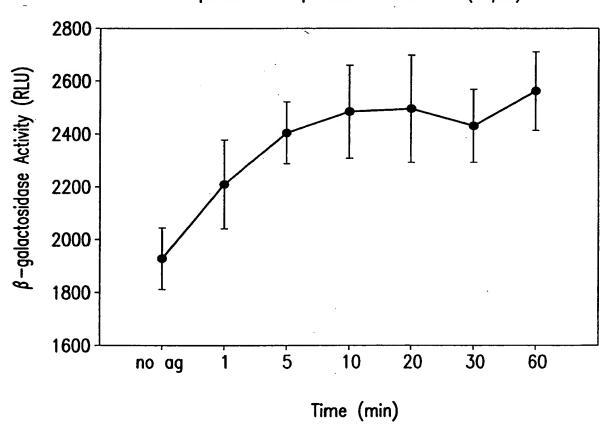


FIG.8B



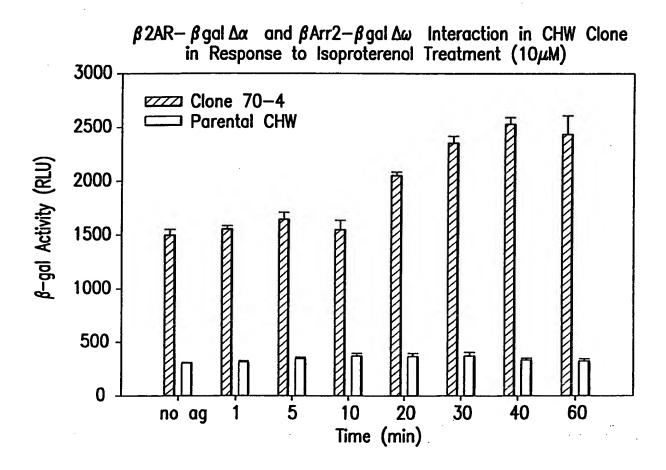


FIG.8C



 β —galactosidase Complementation as a Measurement for Adrenergic Receptor Homodimerization in HEK 293 Cells Coexpressing β 2AR- β gal $\Delta\alpha$ and β 2AR- β gal $\Delta\omega$.

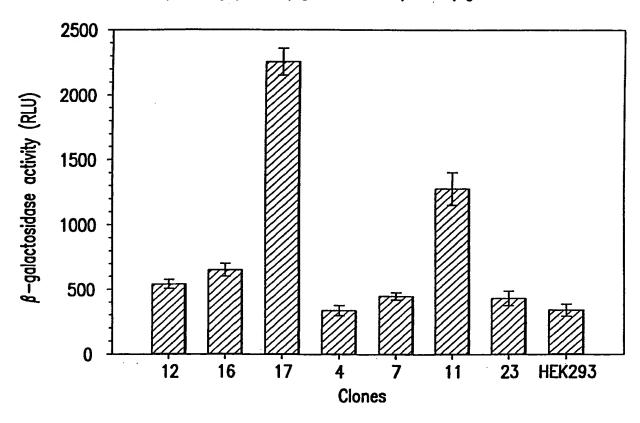


FIG. 9A



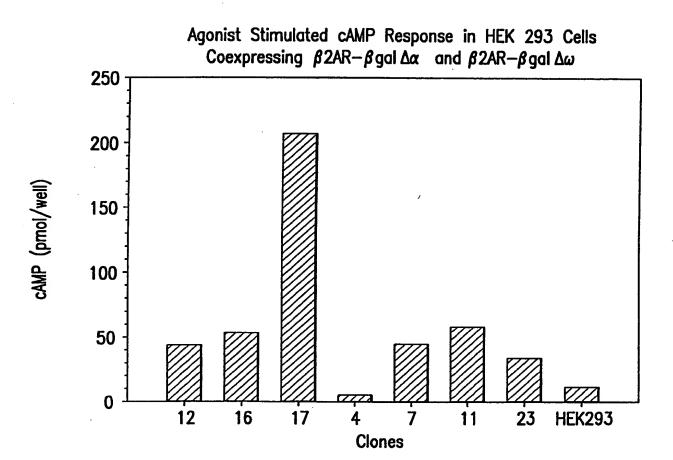


FIG.9B



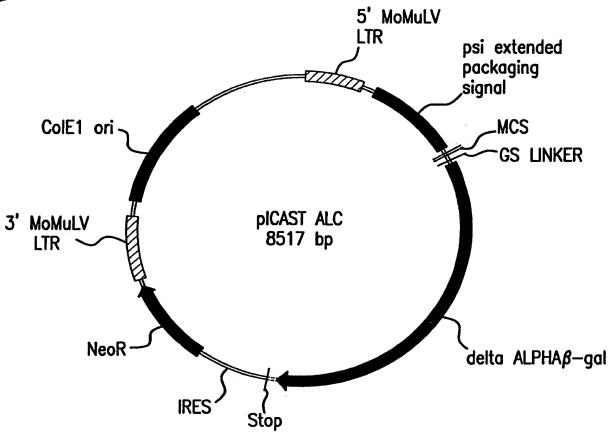


FIG.10A



1		AAACAGGATA TTTGTCCTAT	
51		CAGATGGAAC GTCTACCTTG	
101		TCCTGCCCCG AGGACGGGGC	
151		CCCTCAGCAG GGGAGTCGTC	
201		ACCTGAAATG TGGACTTTAC	
251	 	CGCTTCTGTT GCGAAGACAA	
301		ACCCCTCACT TGGGGAGTGA	
351		CCGTGTATCC GGCACATAGG	
401		TGTTCCTTGG ACAAGGAACC	
451		TCTTTCATTT AGAAAGTAAA	
501		ACCGACCCAC TGGCTGGGTG	
551		CCGATTGTCT GGCTAACAGA	
601		TTAGCTAACT AATCGATTGA	



651		CTGACGAGTT GACTGCTCAA		
701		TTTGGGGGCC AAACCCCCGG		
751		TCCGACCCCG AGGCTGGGGC		
801		AGTTCCCGCC TCAAGGGCGG		
851		GCGTCTTGTC CGCAGAACAG		
901		TTCTGTATTT AAGACATAAA		
951		TTGACCTTAG AACTGGAATC		
1001		GGTAGATGTC CCATCTACAG		
1051		CAACCTTTAA GTTGGAAATT	•	
1101		ATCACCCAGG TAGTGGGTCC		
1151		AGACCAGGTC TCTGGTCCAG		
1201		CTCCCTGGGT GAGGGACCCA		
1251	TCCTCTTCCT AGGAGAAGGA	CCATCCGCCC GGTAGGCGGG		



1301						CCTT											
	GGG	GCG(GAGC	TAC	GGAG	GGA/	A AT	AGG	TCG	iGG	AGT	GAG	GAA	G A	GAT	CCC	GCGG
1351						TTAA1											
	CCG	スしは	AGA I	CGC	אוטנ	WTTA	4 16	CIG	AGI	GΑ	IAI		.GC I	АА	GC I	IAC	11 CC
1401						CCTT AGGAA											
+2		. -	M 1	G	٧.	Ι Τ	· ·	! D	S	L	Α	٧	٧	A	R	. 1	D
1451		GAG/	ATGG	GCG	STG/		GG	ATT	CAC	TG	GCC	GTC	GTG	G C	CCG	CAC	CGA
+2	R	P	S	Q	Q	L	R	S	L	N	G	ì E	: W		R	F	Α
1501						CAATG											
+2	W F	- F	Р А	F) E	Α	٧	P	E		S • • • •	W	L	Ε	С	D	L
1551						AGC6											
+2	P	Ε	A	D	Т	v v	,	V	Р	S	N	W	Q	М	Н	Œ	i Y
1601						TCGT AGCA											
+2	D	Α	Р	I	Y	Т	N	٧	T	Υ	Р	· I	T	,	V	N	Р
1651						ACCA TGGT											
+2	P F	- V	/ P	T	Ε	. N	Р	T	G	(Υ	S	L	T .	F	N
1701	CGTT																



+2	٧])	E	S	h 	I 	L 	Q	E	<u> </u>	G	Q	T	-	R	I	.]	[F	D	G
1751																-						TGG ACC
+2	1	V 	N 	S 		A 	F 	H 	L	-	W	C	N 	- 	G 	R		i 	۷		à '	Y
1801										TGT ACA												
+2	G	Q	D) S	;	R	L 	P	, 	S	E		=	D 	L 		S . 	A		F 	L	R
1851										TCT AGA												
+2	Α		à 	E 	N .	R		L 	Α	۷	' 	M	٧	L	. 1	₹	W		S 	D	G	S
1901										GGT CCA												
+2		/ 	L	Ε		D 	Q 	D 	M	 -`	W	R	. M		S 	G	I		F	F	})
1951										TGT ACA												
+2	٧	S	L	L		Н	K 	P	, 	T	T)	I 	S)	F 		H 	٧	Α
2001										ACT TGA												
+2	T	R	≀	F 	N	D	ا ۔ ۔	D 	F 	S 		₹	Α	V 		- 	Ε		١	E	٧	Q
2051	ACT									CAG GTC												

FIG.10E



+2	M C G E L R D Y L R V T V S L I	W
2101	GATGTGCGGC GAGTTGCGTG ACTACCTACG GGTAACAGTT TCTTTATCCTACACGCCG CTCAACGCAC TGATGGATGC CCATTGTCAA AGAAATAC	
+2	Q G E T Q V A S G T A P F G G E	I
2151	AGGGTGAAAC GCAGGTCGCC AGCGGCACCG CGCCTTTCGG CGGTGAAATCCCACTTTG CGTCCAGCGG TCGCCGTGGC GCGGAAAGCC GCCACTT	
+2	I D E R G G Y A D R V T L R L N	V
2201	ATCGATGAGC GTGGTGGTTA TGCCGATCGC GTCACACTAC GTCTGAAGTAGCTACTCG CACCACCAAT ACGGCTAGCG CAGTGTGATG CAGACTTG	
+2	ENPKLWS AEI PNL YRA	4
2251	CGAAAACCCG AAACTGTGGA GCGCCGAAAT CCCGAATCTC TATCGTGGGCTTTTGGGC TTTGACACCT CGCGGCTTTA GGGCTTAGAG ATAGCACG	
+2	V V E L H T A D G T L I E A E A	C
2301	TGGTTGAACT GCACACCGCC GACGGCACGC TGATTGAAGC AGAAGCCTACCAACTTGA CGTGTGGCGG CTGCCGTGCG ACTAACTTCG TCTTCGGA	
+2	D V G F R E V R I E N G L L L L	N
2351	GATGTCGGTT TCCGCGAGGT GCGGATTGAA AATGGTCTGC TGCTGCTCCTACAGCCAA AGGCGCTCCA CGCCTAACTT TTACCAGACG ACGACGAC	
+2	G K P L L I R G V N R H E H H I)
2401	CGGCAAGCCG TTGCTGATTC GAGGCGTTAA CCGTCACGAG CATCATCG GCCGTTCGGC AACGACTAAG CTCCGCAATT GGCAGTGCTC GTAGTAGG	

FIG.10F



+2	L	Н	G	Q	,	V 1	M -	D	Ε	Q	7	Г I	М	٧	Q	D		Ι	L	L
2451													TGO							
+2	М	K	(ן ן	N	N	F	N	ļ	۱ ۱	V	R	С	S	Н	۱ '	Υ .	Р	N	Н
2501													TGT ACA							
+2	F		L	W	Υ	Т	L	_ (D	R	Υ	(a	L	Υ	٧	٧	. [)
2551													CG(
+2	Ε	A	N	I		Ξ	Γ	Н	G	М	۷	' I	P	М	N	R		L 	T	D
2601													CA/ GTT							
+2	D	P	F	R 1	N	L	P	Α	٨	1 :	S	Ε	R	٧	T	· [₹	М	٧	Q
2651													CGC							
+2	R	\ 	D 	R	N	Н	P		S	۷.	I	I	V	1 :	S ·	L	G	N	E	
2701													CTO GAO							
+2	S	G	Н	G	. /	۱ ۱	1	H .	D	Α	L	. \	Y	R	W .	Ι	ŀ	(S	٧
2751													ATC TAG				-			

FIG.10G



+2	D	Р	S	R	P	V Q	Y	′ E	G	G	G	Α	D	T	Τ	Α
2801				GCC GCG												
+2	T	D	I	I	С	Р	M 	Υ	A F	۱ .	/ D	Ε	D	Q	P	
2851				ATT AAT												
+2	F	· /	A V	Р	K	W	S 	Ι	K	Κ	W	L S	5 l	_ F) (. 	G
2901				GCC												
+2	E	T	R	P	L 	I L		: E	Y	Α	Н	Α	M	G	N	S
2951				CGC GCG												
+2	L	G	G	F	Α	K	Y 	W 	Q A	\ F	R	Q	Υ	Р	R	
3001				TTC AAG												
+2	L () (G G	i F	٧	W	D	W	٧	D	Q	S L	.]	[K	(' 	Υ .
3051				CTT GAA												
+2	D	Ε	N 	G 	N	P W	S 	. A	Υ	G	G 	D	F	G	D	T
3101				GCA CGT												

FIG.10H



+2	Р	N	D	R	Q	F	С	М	N	G	L	٧	F	Α	D	R
3151											TCTC					
+2	T	P F	l P	Α	L	T	E	Α	K		1 0	Q	Q	F	F	Q
3201											ACCA TGGT					
+2	F	R	L :	S (G (Q T	·	[[E V	/	Т	S	E '	Y 	L I	F R
3251											ACCA TGGT					
+2	. Н	S	D	N	Ε	L	L 	Н	W	M	٧	Α	L	D	G	K
3301											GGTG CCAC					
+2	P L	_ A	S	G	Ε	V	P	L	D	٧	' A	. P	Q	G	K	Q
3351											TCGC AGCG					
+2	L 	Ι	E I	L 6) [E L		·) P	, 	E	S · /	A (à (Q I	. W
3401											GAGA CTCT				_	
+2	L	T	٧	R	٧	٧	Q 	Р 	N 	A 	T	Α	W	S	E	Α
3451											GACC CTGG					

FIG.101



+2	G 	Н	I	S 		A !	W	Q	Q	W	F	? 1	L	Α	Ε	N		L 	S	٧
3501									CAG											
+2	T	L		P ,	A 	Α	S	Н	Α	. :	Ι	Р	Н	L	T		T	S	Ε	М
3551									CGC							-				
+2		D	F	C	I	Ε	L	(3	N 	K	R		d (Q 	F 	N	R	. (Q
3601									GTA CAT											
+2	S 	G	F 	L		S (Q 	M 	W	I	9	i [)	K	K	Q	ا 	_	L	T
3651									TGG ACC								-			
+2	Р	L	. I	R 1	D	Q	F	T	R		۹	P	L	D	N	[D 	I	G	٧
3701									CCG								_			
+2		S 	E	Α	Т	R	I) 	P	N	Α	, , , ,	۷ ۱	V	Ε	R	W	' 	(
3751									ACC TGG											
+2	Α	Α	G 	Н		· (Q ,	A 	Ε	Α	Α	. l	- 	L	Q	C	7	Γ 	Α	D
3801									GAA CTT											

FIG.10J



+2	T 	L	Α	D	Α	V L	. 1		Т	· • •	Α	H 	Α	W	Q	Н	Q
3851						TGCT ACGA											
+2	G	. K	T	L	F	I	S	R	K	T 	Υ	R	I) (i S	3
3901						TATCA TAGT											
+2	G (Q	М ,	\]	. 1	· V	D	٧	Ε	۷		A :	S [)	T	P	Н
3951						CGTT											
+2	Р	A	R	I	G	L N	C) L		Α	Q	٧	Α	Ε	R	٧
4001						TGAA ACTT											
+2	N	W	L	G	L	G	Р	Q	E	N	Υ	Р	D	R	L	. 1	•
4051						GGGC CCCG											
+2	A /	A (C F		R	W	D	L	Р	L		5	D 1	1	Υ	T	Р
4101						CTGG GACC											
+2	Т	۷	F	P	S	E N	G	i L	. R		С	G	T	R	Ε	L	N
4151						AAAA											

FIG.10K



+2	,	Y	G	P	H	1 (Q	W	R		G	D	F	:	Q	F	:	N	I		S	R
4201																						GCT GCGA
+2	Υ	S	Q	0) 	Q	L 	М		Ε	T		S	H 		R 	Н	L		L	Н	Α
4251											-											GCG GCGC
+2	Ε		Ξ	G 	T 	W 		_	١	I 	·	D 	G 		F	H 	M	1 1	G 	I		G G
4301																						ACC
+2) 	D	S 	h	l :	S 	Р	S		۷	S 	Α		Ε	F	:	Q	L		S 	Α
4351																			_			GGC GGC
+2	G	R	Υ	Н		Υ	Q	L		٧	W		C 	Q 		(R	S 		D 	Υ	K
4401																						TAAA NTTT
+2	D		Ξ	D 	L 	D		1 H	∤ • • •	Н	 	H 	H 		H 	R · - >						
4451					_									-								AGA ATCT
4501												-	_		-							TTC AAG
4551							-															CTG GAC

FIG.10L



pICAST ALC

4601			CCCCTCTCGC GGGGAGAGCG	
4651			GTTCCTCTGG CAAGGAGACC	
4701			CAGGCAGCGG GTCCGTCGCC	
4751	 		CACGTGTATA GTGCACATAT	
4801			GTGAGTTGGA CACTCAACCT	
4851			CAACAAGGGG GTTGTTCCCC	
4901			ATCTGGGGCC TAGACCCCGG	
4951	_		AACGTCTAGG TTGCAGATCC	
5001		= 1 1	CGATGATAAT GCTACTATTA	
5051			CCGCTTGGGT GGCGAACCCA	
5101			GGCTGCTCTG CCGACGAGAC	
,5151			TCTTTTTGTC AGAAAAACAG	

FIG.10M



pICAST ALC

5201			CTGCAGGACG		
	ACAGGCCACG	GGACTTACTT	GACGTCCTGC	TCCGTCGCGC	CGATAGCACC
5251	CTGGCCACGA	CGGGCGTTCC	TTGCGCAGCT	GTGCTCGACG	TTGTCACTGA
	GACCGGTGCT	GCCCGCAAGG	AACGCGTCGA	CACGAGCTGC	AACAGTGACT
5301			TATTGGGCGA		
-			ATAACCCGCT		
5351			GCCGAGAAAG		
			CGGCTCTTTC		,
5401			TGATCCGGCT		
			ACTAGGCCGA		
5451			GAGCACGTAC		
			CTCGTGCATG		
5501			GAAGAGCATC		
			CTTCTCGTAG	-	
5551			GCGCATGCCC		
= = = =			CGCGTACGGG		•
5601	· · · · ·		TGCCGAATAT		
			ACGGCTTATA		
5651			GGCCGGCTGG		
			CCGGCCGACC		
5701					GCGGCGAATG
			ACTATAACGA		•
5751			TTTACGGTAT		
	CCGACTGGCG	AAGGAGCACG	AAATGCCATA	GCGGCGAGGG	CTAAGCGTCG

FIG.10N



pICAST ALC

5801		CTATCGCCTT GATAGCGGAA	 	
5851	-	GATAAAATAA CTATTTTATT	 	
5901		ACCCCACCTG TGGGGTGGAC		
5951		GCATGGAAAA CGTACCTTTT		
6001		AACAGATGGA TTGTCTACCT		
6051		GTTCCTGCCC CAAGGACGGG		
6101		CCAAACAGGA GGTTTGTCCT	 	
6151		AACAGATGGT TTGTCTACCA	 	
6201		ATCAGATGTT TAGTCTACAA	 	
6251		TTTGAACTAA AAACTTGATT		
6301		TCCCCGAGCT AGGGGCTCGA		
6351		CTCCGATTGA GAGGCTAACT		

FIG.100



pICAST ALC

6401		CCGACTTGTG GGCTGAACAC		
6451		CTACCCGTCA GATGGGCAGT		
6501		TTGGTTTTT AACCAAAAAA	•	
6551		GAATCGGCCA CTTAGCCGGT		
6601		CTTCCTCGCT GAAGGAGCGA		
6651		GTATCAGCTC CATAGTCGAG		

FIG.10P



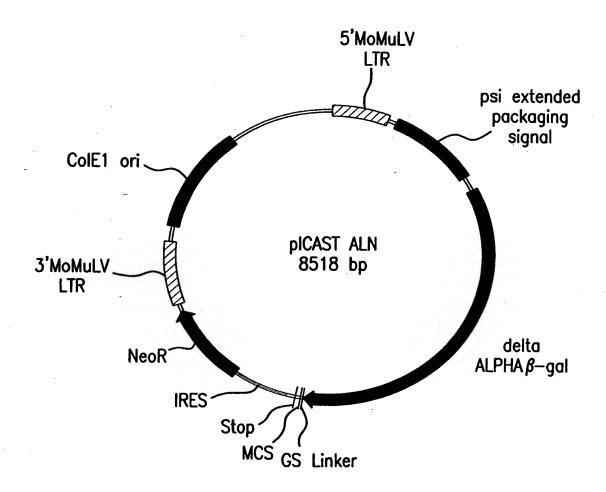


FIG.11A



pICAST ALN

CTGCAGCCTG	AATATGGGCC	AAACAGGATA	TCTGTGGTAA	GCAGTTCCTG	CCCCGGCTCA	60
GACGTCGGAC	TTATACCCGG	TTTGTCCTAT	AGACACCATT	CGTCAAGGAC	GGGGCCGAGT	60
GGGCCAAGAA	CAGATGGAAC	AGCTGAATAT	GGGCCAAACA	GGATATCTGT	GGTAAGCAGT	120
CCCGGTTCTT	GTCTACCTTG	TCGACTTATA	CCCGGTTTGT	CCTATAGACA	CCATTCGTCA	120
	GCTCAGGGCC					180
	CGAGTCCCGG					180
	ACCATCAGAT					240
	TGGTAGTCTA					240
	TAACCAATCA					300
	ATTGGTTAGT					300
	AGAGCCCACA					360
	TCTCGGGTGT					360
	CCGTGTATCC					420
	GGCACATAGG					420
	GAGGGTCTCC					480
	CTCCCAGAGG					480
	CCGGGATCGG					540
	GGCCCTAGCC					540
	AGCAACTTAT		· · · · -			600
	TCGTTGAATA					600
	TCGGTACTAG					660
	AGCCATGATC					660
	CTGAACACCC					720.
	GACTTGTGGG					720
	CCCGACCTGA					780
CAAAAACACC	GGGCTGGACT	CCTTCCCTCA	GUTACACCTT	AGGC I GGGGC	AGICCIATAC	780



pICAST ALN

	AGGAGACGAG					840
ACCAAGACCA	TCCTCTGCTC	TTGGATTTTG	TCAAGGGCGG	AGGCAGACTT	AAAAACGAAA	840
CGGTTTGGAA	CCGAAGCCGC	GCGTCTTGTC	TGCTGCAGCA	TCGTTCTGTG	TTGTCTCTGT	900
	GGCTTCGGCG					900
_	TTCTGTATTT					960
	AAGACATAAA					960
	GTAACTGGAA					1020
	CATTGACCTT					1020
	GTTGGGTTAC					1080
	CAACCCAATG					1080
	GCACCTTTAA					1140
	CGTGGAAATT					1140
	ATGGACACCC					1200
	TACCTGTGGG					1200
	CTCCCTGGGT					1260
	GAGGGACCCA					1260
	CGTCTCTCCC					1320
	GCAGAGAGGG					1320
	TCACTCCTTC					1380
	AGTGAGGAAG					1380
					AAGATGAGGA	1440
	AGCTTGTGGT					1440
	GGCGTGATTA					1500
	CCGCACTAAT					1500
	CGCAGCCTGA					1560
GGTTGTCAAT	GCGTCGGACT	TACCGCTTAC	CGCGAAACGG	ACCAAAGGCC	GIGGICTTCG	1560



pICAST ALN

GGTGCCGGAA	AGCTGGCTGG	AGTGCGATCT	TCCTGAGGCC	GATACTGTCG	TCGTCCCCTC	1620
CCACGGCCTT	TCGACCGACC	TCACGCTAGA	AGGACTCCGG	CTATGACAGC	AGCAGGGGAG	1620
AAACTGGCAG	ATGCACGGTT	ACGATGCGCC	CATCTACACC	AACGTGACCT	ATCCCATTAC	1680
	TACGTGCCAA					1680
	CCGTTTGTTC					1740
	GGCAAACAAG					1740
	AGCTGGCTAC					1800
	TCGACCGATG					1800
	CTGTGGTGCA					1860
	GACACCACGT					1860
	GACCTGAGCG					1920
	CTGGACTCGC					1920
	AGTGACGGCA					1980
	TCACTGCCGT					1980
	GTCTCGTTGC					2040
	CAGAGCAACG					2040
	AATGATGATT					2100
	TTACTACTAA					2100
	GACTACCTAC					2160
	CTGATGGATG					2160
	GCGCCTTTCG					2220
	CGCGGAAAGC					2220
	CGTCTGAACG					2280
	GCAGACTTGC					2280
	GTGGTTGAAC					2340
GATAGCACGC	CACCAACTTG	ACGIGTGGCG	GCTGCCGTGC	GACTAACTTC	GTCTTCGGAC	2340



CGATGTCGGT	TTCCGCGAGG	TGCGGATTGA	AAATGGTCTG	CTGCTGCTGA	ACGGCAAGCC	2400
	AAGGCGCTCC					2400
GTTGCTGATT	CGAGGCGTTA	ACCGTCACGA	GCATCATCCT	CTGCATGGTC	AGGTCATGGA	2460
CAACGACTAA	GCTCCGCAAT	TGGCAGTGCT	CGTAGTAGGA	GACGTACCAG	TCCAGTACCT	2460
	ATGGTGCAGG					2520
ACTCGTCTGC	TACCACGTCC	TATAGGACGA	CTACTTCGTC	TTGTTGAAAT	TGCGGCACGC	2520
	TATCCGAACC					2580
GACAAGCGTA	ATAGGCTTGG	TAGGCGACAC	CATGTGCGAC	ACGCTGGCGA	TGCCGGACAT	2580
	GAAGCCAATA			, –		2640
ACACCACCTA	CTTCGGTTAT	AACTTTGGGT	GCCGTACCAC	GGTTACTTAG	CAGACTGGCT	2640
	TGGCTACCGG					2700
ACTAGGCGCG	ACCGATGGCC	GCTACTCGCT	TGCGCATTGC	GCTTACCACG	TCGCGCTAGC	2700
TAATCACCCG	AGTGTGATCA	TCTGGTCGCT	GGGGAATGAA	TCAGGCCACG	GCGCTAATCA	2760
ATTAGTGGGC	TCACACTAGT	AGACCAGCGA	CCCCTTACTT	AGTCCGGTGC	CGCGATTAGT	2760
CGACGCGCTG	TATCGCTGGA	TCAAATCTGT	CGATCCTTCC	CGCCCGGTGC	AGTATGAAGG	2820
GCTGCGCGAC	ATAGCGACCT	AGTTTAGACA	GCTAGGAAGG	GCGGGCCACG	TCATACTTCC	2820
CGGCGGAGCC	GACACCACGG	CCACCGATAT	TATTTGCCCG	ATGTACGCGC	GCGTGGATGA	2880
GCCGCCTCGG	CTGTGGTGCC	GGTGGCTATA	ATAAACGGGC	TACATGCGCG	CGCACCTACT	2880
AGACCAGCCC	TTCCCGGCTG	TGCCGAAATG	GTCCATCAAA	AAATGGCTTT	CGCTACCTGG	2940
TCTGGTCGGG	AAGGCCGAC	ACGGCTTTAC	CAGGTAGTTT	TTTACCGAAA	GCGATGGACC	2940
	CCGCTGATCC					3000
TCTCTGCGCG	GGCGACTAGG	AAACGCTTAT	GCGGGTGCGC	TACCCATTGT	CAGAACCGCC	3000
TTTCGCTAAA	TACTGGCAGG	CGTTTCGTCA	GTATCCCCGT	TTACAGGGCG	GCTTCGTCTG	3060
AAAGCGATTT	ATGACCGTCC	GCAAAGCAGT	CATAGGGGCA	AATGTCCCGC	CGAAGCAGAC	3060
GGACTGGGTG	GATCAGTCGC	TGATTAAATA	TGATGAAAAC	GGCAACCCGT	GGTCGGCTTA	3120
CCTGACCCAC	CTAGTCAGCG	ACTAATTTAT	ACTACTTTTG	CCGTTGGGCA	CCAGCCGAAT	3120



	TTTGGCGATA AAACCGCTAT					3180 3180
	ACGCCGCATC TGCGGCGTAG					3240 3240
	TCCGGGCAAA AGGCCCGTTT					3300 3300
	CTGCACTGGA GACGTGACCT					3360 3360
	GTCGCTCCAC CAGCGAGGTG					3420 3420
	GGGCAACTCT CCCGTTGAGA					3480 3480
	GGGCACATCA CCCGTGTAGT					3540 3540
	GCCGCGTCCC CGGCGCAGGG					3600 3600
	GGTAATAAGC CCATTATTCG					3660 3660
	GATAAAAAAC CTATTTTTG					3720 3720
					CCTGGGTCGA GGACCCAGCT	3780 3780
TGCGACCTTC	CGCCGCCCGG	TAATGGTCCG	GCTTCGTCGC	AACAACGTCA		3840 3840
	GATGCGGTGC CTACGCCACG				AGGGGAAAAC TCCCCTTTTG	3900 3900



AGCCGGAAAA TCGGCCTTTT			3960 3960
GTGGCGAGCG CACCGCTCGC			4020 4020
GCAGAGCGGG CGTCTCGCCC			4080 4080
GCCGCCTGTT CGGCGGACAA			4140 4140
CCGAGCGAAA GGCTCGCTTT			4200 4200
CGCGGCGACT GCGCCGCTGA			4260 4260
CATCGCCATC GTAGCGGTAG			4320 4320
GGGATTGGTG CCCTAACCAC			4380 4380
GGTCGCTACC CCAGCGATGG			4440 4440
CCTTGGCGCG GGAACCGCGC			4500 4500
GATTAGATGC CTAATCTACG			4560 4560
GGCAATGTGA CCGTTACACT			462 0 462 0
TCCCCTCTCG AGGGGAGAGC			4680 4680



GAAGCTTCTT CTTCGAAGAA			4740 4740
CCTGGCGACA GGACCGCTGT			4800 4800
GCACAACCCC CGTGTTGGGG			4860 4860
TCAAGCGTAT AGTTCGCATA			4920 4920
GATCTGGGGC CTAGACCCCG			4980 4980
GCCCCCGAA CGGGGGGCTT			5040 5040
GAACAAGATG CTTGTTCTAC			5100 5100
GACTGGGCAC CTGACCCGTG			5160 5160
GGGCGCCCGG CCCGCGGGCC			5220 5220
GAGGCAGCGC CTCCGTCGCG			5280 5280
GTTGTCACTG CAACAGTGAC		AAGTGCCGGG TTCACGGCCC	5340 5340
CTGTCATCTC GACAGTAGAG			5400 5400
CTGCATACGC GACGTATGCG		AAGCGAAACA TTCGCTTTGT	5460 5460



TCGCATCGAG	CGAGCACGTA	CTCGGATGGA	AGCCGGTCTT	GTCGATCAGG	ATGATCTGGA	5520
AGCGTAGCTC	GCTCGTGCAT	GAGCCTACCT	TCGGCCAGAA	CAGCTAGTCC	TACTAGACCT	5520
CGAAGAGCAT	CAGGGGCTCG	CGCCAGCCGA	ACTGTTCGCC	AGGCTCAAGG	CGCGCATGCC	5580
GCTTCTCGTA	GTCCCCGAGC	GCGGTCGGCT	TGACAAGCGG	TCCGAGTTCC	GCGCGTACGG	5580
				TTGCCGAATA		5640
-				AACGGCTTAT		5640
				GGTGTGGCGG		5700
				CCACACCGCC		5700
				GGCGGCGAAT		5760
	•			CCGCCGCTTA		5760
				CGCATCGCCT		5820
				GCGTAGCGGA		5820
				CGATAAAATA		5880
AGAACTGCTC	AAGAAGACTC	GCCCTGAGAC	CCCAAGCGTA	GCTATTTTAT	TTTCTAAAAT	5880
				GTAGGTTTGG		5940
				CATCCAAACC		5940
				TGAGAATAGA		6000
				ACTCTTATCT		6000
			•	ACAGGATATC		6060
. ,				TGTCCTATAG		6060
				CTGAATATGG		6120
				GACTTATACC		6120
				GAACAGATGG		6180
				CTTGTCTACC		6180
				TTCCAGGGTG		6240
GCCAGGTCGG	GAGTCGTCAA	AGATCTCTTG	GTAGTCTACA	AAGGTCCCAC	GGGGTTCCTG	6240



CCTGTGCCTT GGACACGGAA				6300 6300
CTCCCCGAGC GAGGGGCTCG				6360 6360
ACTGAGTCGC TGACTCAGCG		 		6420 6420
 GGTCTCGCTG CCAGAGCGAC		 		6480 6480
TTTCATTCAT AAAGTAAGTA				6540 6540
AATGGCCATA TTACCGGTAT				6600 6600
 CGCGAGAAGG GCGCTCTTCC		 		6660 6660
GGTATCAGCT CCATAGTCGA				67 20 67 20
AAAGAACATG TTTCTTGTAC	· · · - · - · -	 		6780 6780
GGCGTTTTTC CCGCAAAAAG				6840 6840
GAGGTGGCGA CTCCACCGCT			_	6900 6900
 		 	CTGTCCGCCT GACAGGCGGA	69 60 69 60
GGGAAGCGTG CCCTTCGCAC				7020 7020



TGTAGGTCGT	TCGCTCCAAG	CTGGGCTGTG	TGCACGAACC	CCCCGTTCAG	CCCGACCGCT	7080
ACATCCAGCA	AGCGAGGTTC	GACCCGACAC	ACGTGCTTGG	GGGGCAAGTC	GGGCTGGCGA	7080
GCGCCTTATC	CGGTAACTAT	CGTCTTGAGT	CCAACCCGGT	AAGACACGAC	TTATCGCCAC	7140
CGCGGAATAG	GCCATTGATA	GCAGAACTCA	GGTTGGGCCA	TTCTGTGCTG	AATAGCGGTG	7140
TGGCAGCAGC	CACTGGTAAC	AGGATTAGCA	GAGCGAGGTA	TGTAGGCGGT	GCTACAGAGT	7200
-	GTGACCATTG					7200
	GTGGCCTAAC					7260
•	CACCGGATTG					7260
	AGTTACCTTC					7320
	TCAATGGAAG					7320
	CGGTGGTTTT					7380
	GCCACCAAAA					7380
	TCCTTTGATC					7440
	AGGAAACTAG					7440
	TTTGGTCATG					7500
	AAACCAGTAC					7500
	TCTAAAGTAT					7560
	AGATTTCATA					7560
	CTATCTCAGC			· · · · · · · · · · · · · · · · · ·		7620
	GATAGAGTCG					7620
					TGCAATGATA	7680
	ATTGATGCTA		•			7680
	CACGCTCACC					7740
	GTGCGAGTGG					7740
	GAAGTGGTCC					7800
CGGCTCGCGT	CTTCACCAGG	ACGTTGAAAT	AGGCGGAGGT	AGGTCAGATA	ATTAACAACG	7800



CGGGAAGCTA GAGTAAGTAGGCCCTTCGAT CTCATTCATC			 	7860 7860
ACAGGCATCG TGGTGTCACG				792 0 792 0
CGATCAAGGC GAGTTACATC				7980 7980
CCTCCGATCG TTGTCAGAAGGGAGGCTAGC AACAGTCTTG			 	804 0 804 0
CTGCATAATT CTCTTACTGT GACGTATTAA GAGAATGACA	·			8100 8100
TCAACCAAGT CATTCTGAGAAGTTGGTTCA GTAAGACTCT			 	8160 8160
ATACGGGATA ATACCGCGCCCTAT TATGGCGCGCGC				8220 8220
TCTTCGGGGC GAAAACTCTC			 	8280 8280
ACTCGTGCAC CCAACTGATO		_	 	8340 8340
AAAACAGGAA GGCAAAATGC				8400 8400
CTCATACTCT TCCTTTTTCA GAGTATGAGA AGGAAAAAGT			 	8460 8460
GGATACATAT TTGAATGTAT				8518 8518

FIG.11L



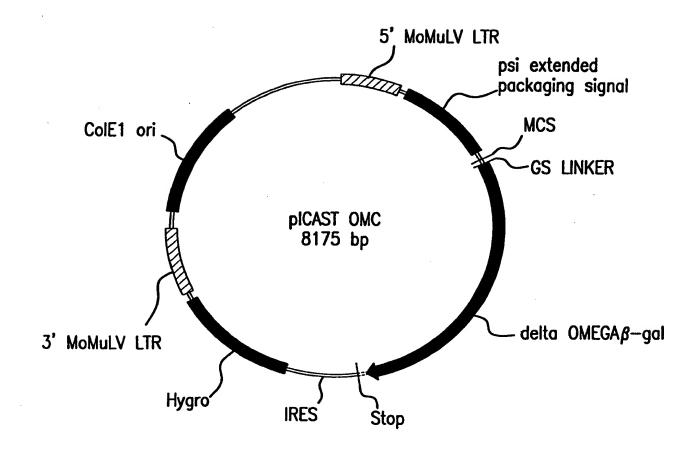


FIG.12A



		TCTGTGGTAA AGACACCATT			60 60
		GGGCCAAACA CCCGGTTTGT			120 120
		GGTCCCCAGA CCAGGGGTCT			180 180
		TGCCCCAAGG ACGGGGTTCC			240 240
		CGCTTCTGTT GCGAAGACAA			300 300
		CGGGGCGCCA GCCCCGCGGT			360 360
	- -	CTTGCAGTTG GAACGTCAAC			420 420
		TGACTACCCG ACTGATGGGC			480 480
		CCCAGGGACC GGGTCCCTGG			540 540
		CCGATTGTCT GGCTAACAGA	- ·		600 600
		AGCTCTGTAT TCGAGACATA		CGTGGTGGAA GCACCACCTT	660 660
		CTGGGAGACG GACCCTCTGC			720 720
		CGATGTGGAA GCTACACCTT		•	780 780



TGGTTCTGGT	AGGAGACGAG	AACCTAAAAC	AGTTCCCGCC	TCCGTCTGAA	ТППССПТ	840
	TCCTCTGCTC					840
	CCGAAGCCGC					900
GCCAAACCTT	GGCTTCGGCG	CGCAGAACAG	ACGACGTCGT	AGCAAGACAC	AACAGAGACA	900
CTGACTGTGT	TTCTGTATTT	CTCTCAAAAT	TACCCCCACA	CTCTTACCAC	TCCCTTAACT	060
	AAGACATAAA					96 0 96 0
		Charottin	Aiccodaici	anomitatia	AdddAATTCA	300
TTGACCTTAG	GTAACTGGAA	AGATGTCGAG	CGGCTCGCTC	ACAACCAGTC	GGTAGATGTC	1020
AACTGGAATC	CATTGACCTT	TCTACAGCTC	GCCGAGCGAG	TGTTGGTCAG	CCATCTACAG	1020

	GTTGGGTTAC					1080
TICTICICIG	CAACCCAATG	GAAGACGAGA	CGICTIACCG	GIIGGAAAII	GCAGCCTACC	1080
CCGCGAGACG	GCACCTTTAA	CCGAGACCTC	ATCACCCAGG	TTAAGATCAA	GGTCTTTTCA	1140
	CGTGGAAATT					1140
	ATGGACACCC					1200
GGACCGGGCG	TACCTGTGGG	TCTGGTCCAG	GGGATGTAGC	ACTGGACCCT	TCGGAACCGA	1200
TTTCACCCCC	CTCCCTCCCT	CAACCCCTTT	CTACACCCTA	ACCCTCCCCC	TOOTOTTOOT	1000
	CTCCCTGGGT GAGGGACCCA					1260 1260
7 V V V V V V V V V V V V V V V V V V V	anddancech	a i i cada	CATGIGGGAI	rcddAddcdd	AGGAGAAGGA	1200
CCATCCGCCC	CGTCTCTCCC	CCTTGAACCT	CCTCGTTCGA	CCCCGCCTCG	ATCCTCCCTT	1320
GGTAGGCGGG	GCAGAGAGGG	GGAACTTGGA	GGAGCAAGCT	GGGGCGGAGC	TAGGAGGGAA	1320
	TCACTCCTTC					1380
ATAGGTCGGG	AGTGAGGAAG	AGATCCGCGG	CCGGCGAGAT	CGGGTAATTA	TGCTGAGTGA	1380
ATAGGGCGAT	TCGAATCAGG	CCTTGGCGCG	CCGGATCCTT	AATTAAGCGC	AATTGGGAGG	1440
	AGCTTAGTCC					1440
		44 1.004040			11701000100	1110
TGGCGGTAGC	CTCGAGATGG	GCGTGATTAC	GGATTCACTG	GCCGTCGTTT	TACAACGTCG	1500
ACCGCCATCG	GAGCTCTACC	CGCACTAATG	CCTAAGTGAC	CGGCAGCAAA	ATGTTGCAGC	1500
TOACTOOOAA	110007000		T T	00400		
	AACCCTGGCG					1560
ACTUACCCIT	TTGGGACCGC	AATGGGTTGA	ATTAGUGGAA	CUICUIGIAG	GGGGAAAGCG	1560

FIG.12C



AATAGCGAAG TTATCGCTTC		 	1620 1620
TGGCGCTTTG ACCGCGAAAC			1680 1680
CTTCCTGAGG GAAGGACTCC			1740 1740
CCCATCTACA GGGTAGATGT			1800 1800
AATCCGACGG TTAGGCTGCC			1860 1860
CAGACGCGAA GTCTGCGCTT		 	1920 1920
TGGGTCGGTT ACCCAGCCAA		 	1980 1980
CGCGCCGGAG GCGCGGCCTC			2040 2040
GAAGATCAGG CTTCTAGTCC		 	2100 2100
CCGACTACAC GGCTGATGTG			2160 2160
		 GTGACTACCT CACTGATGGA	2220 2220
GTTTCTTTAT CAAAGAAATA			2280 2280
ATTATCGATG TAATAGCTAC		 TACGTCTGAA ATGCAGACTT	2340 2340



	CCGAAACTGT GGCTTTGACA					2400 2400
	GCCGACGGCA CGGCTGCCGT					2460 2460
	GAAAATGGTC CTTTTACCAG					2520 2520
	GAGCATCATC CTCGTAGTAG					2580 2580
	CTGATGAAGC GACTACTTCG					2640 2640
	TGGTACACGC ACCATGTGCG					2700 2700
	CACGGCATGG GTGCCGTACC					2760 2760
	GAACGCGTAA CTTGCGCATT					2820 2820
	CTGGGGAATG GACCCCTTAC					2880 2880
	GTCGATCCTT CAGCTAGGAA					2940 2940
	ATTATTTGCC TAATAAACGG					3000 3000
ACACGGCTTT	ACCAGGTAGT	TTTTTACCGA	AAGCGATGGA	CCTCTCTGCG		3060 3060
					AATACTGGCA TTATGACCGT	



		CAGTATCCCC GTCATAGGGG					3180 3180
		TATGATGAAA ATACTACTTT	· ·				3240 3240
		GATCGCCAGT CTAGCGGTCA			· ·		3300 3300
		ACGGAAGCAA TGCCTTCGTT					3360 3360
		GTGACCAGCG CACTGGTCGC					3420 3420
		CTGGATGGTA GACCTACCAT					3480 3480
		CAGTTGATTG GTCAACTAAC					3540 3540
		GTACGCGTAG CATGCGCATC					3600 3600
(STCGCGGACC	CAGCAGTGGC GTCGTCACCG	CAGACCGCCT	TTTGGAGTCA	CACTGCGAGG	GGCGCCCAG	3660 3660
(GTGCGGTAG	CCGCATCTGA GGCGTAGACT	GGTGGTCGCT	TTACCTAAAA	ACGTAGCTCG	ACCCATTATT	3720 3720
(CGCAACCGTT	AAATTGGCGG	TCAGTCCGAA	AGAAAGTGTC	TACACCTAAC		3780 3780
1	GTTGACGAC	ACGCCGCTGC TGCGGCGACG	CGCTAGTCAA	GTGGGCACAG	CTATCTAGAC	TTGTCTTTGA	3840 3840
		GAAGACCTAG CTTCTGGATC				AATAGGTAGA TTATCCATCT	3900 3900



TAAGTGACTG	ATTAGATGCA	TTTCGACTAG	ATCCCTCGAC	CAATTCCGGT	TATTTTCCAC	3960
ATTCACTGAC	TAATCTACGT	AAAGCTGATC	TAGGGAGCTG	GTTAAGGCCA	ATAAAAGGTG	3960
CATATTGCCG	TCTTTTGGCA	ATGTGAGGGC	CCGGAAACCT	GGCCCTGTCT	TCTTGACGAG	4020
GTATAACGGC	AGAAAACCGT	TACACTCCCG	GGCCTTTGGA	CCGGGACAGA	AGAACTGCTC	4020
CATTCCTAGG	GGTCTTTCCC	CTCTCGCCAA	AGGAATGCAA	GGTCTGTTGA	ATGTCGTGAA	4080
	CCAGAAAGGG					4080
	CCTCTGGAAG					4140
•	GGAGACCTTC					4140
	CCCCCACCTG					4200
	GGGGGTGGAC					4200
	AAGGCGGCAC					4260
	TTCCGCCGTG					4260
	CTCTCCTCAA					4320
ï	GAGAGGAGTT					4320
	GGATCTGATC	,				4380
•	CCTAGACTAG					4380
	GTCTAGGCCC					4440
	CAGATCCGGG					4440
	ATGAAAAAGC					4500
	TACTTTTTCG					4500
	AGCGTCTCCG					4560
	TCGCAGAGGC					4560
	GTAGGAGGGC					4620
•	CATCCTCCCG	1				4620
	CGTTATGTTT					4680
GAIGIIICIA	GCAATACAAA	TAGCCGTGAA	ACG LAGCCGG	UGUGAGGCT	AAGGCCTTCA	4680



	GGGGAATTTA CCCCTTAAAT		 		4740 4740
	CAAGACCTGC GTTCTGGACG				4800 4800
	GCGATCGCTG CGCTAGCGAC				4860 4860
	ATCGGTCAAT TAGCCAGTTA				4920 4920
The second secon	CACTGGCAAA GTGACCGTTT				4980 4980
	CTGATGCTTT GACTACGAAA				5040 5040
	TCCAACAATG AGGTTGTTAC				5100 5100
	ATGTTCGGGG TACAAGCCCC		 		5160 5160
	TGTATGGAGC ACATACCTCG				5220 5220
	CGGCTCCGGG GCCGAGGCCC		 		5280 5280
		•		ACGCAATCGT TGCGTTAGCA	5340 5340
	GCCGGGACTG CGGCCCTGAC				5400 5400
	TGTGTAGAAG ACACATCTTC				5460 5460



GAGGGCAAAG GAATAGA CTCCCGTTTC CTTATCT					5520 5520
TTAGTCTCCA GAAAAAG AATCAGAGGT CTTTTTC					5580 5580
AAGTAACGCC ATTTTGC					5640 5640
CAAGGTCAGG AACAGATG GTTCCAGTCC TTGTCTAG					5700 5700
GTTCCTGCCC CGGCTCAC CAAGGACGGG GCCGAGTC					5760 5760
TATCTGTGGT AAGCAGT ATAGACACCA TTCGTCA	•				5820 5820
GGTCCAGCCC TCAGCAGCCCCAGGTCGGG AGTCGTCA			-		5880 5880
TGAAATGACC CTGTGCCACTTACTGG GACACGGA	AAT AAACTTGATT	GGTTAGTCAA	GCGAAGAGCG	AAGACAAGCG	59 40 59 40
GCGCTTCTGC TCCCCGA(CGA GTTATTTTCT	CGGGTGTTGG	GGÁGTGAGCC	CCGCGGTCAG	60 00 60 00
CTCCGATTGA CTGAGTCG	CGG GCCCATGGGC	ACATAGGTTA	TTTGGGAGAA	CGTCAACGTA	60 60 60 60
CCGACTTGTG GTCTCGCT	ACA AGGAACCCTC	CCAGAGGAGA	CTCACTAACT	GATGGGCAGT	6120 6120
GCGGGGGTCT TTCATTCA CGCCCCCAGA AAGTAAGT	TAC GTCGTACATA	GTTTTAATTA	AACCAAAAA	AAGAATTCAT	6180 6180
TTTACATTAA ATGGCCAT			·		6240 6240



TGCGTATTGG CGCTCTTCCG ACGCATAACC GCGAGAAGGC	 			6300 6300
GCGGCGAGCG GTATCAGCTC CGCCGCTCGC CATAGTCGAG				6360 6360
TAACGCAGGA AAGAACATGT ATTGCGTCCT TTCTTGTACA				6420 6420
CGCGTTGCTG GCGTTTTTCC GCGCAACGAC CGCAAAAAGG	•	· -		6480 6480
CTCAAGTCAG AGGTGGCGAA GAGTTCAGTC TCCACCGCTT	*		*	6540 6540
AAGCTCCCTC GTGCGCTCTC TTCGAGGGAG CACGCGAGAG	 			6600 6600
TCTCCCTTCG GGAAGCGTGG AGAGGGAAGC CCTTCGCACC	 			6660 6660
GTAGGTCGTT CGCTCCAAGC CATCCAGCAA GCGAGGTTCG	 			6720 .6720
CGCCTTATCC GGTAACTATC GCGGAATAGG CCATTGATAG	 		· · · · · · · · · · · · · · · · · · ·	6780 6780
GGCAGCAGCC ACTGGTAACA CCGTCGTCGG TGACCATTGT	 			6840 6840
CTTGAAGTGG TGGCCTAACT GAACTTCACC ACCGGATTGA	 			6900 6900
GCTGAAGCCA GTTACCTTCG CGACTTCGGT CAATGGAAGC				6960 6960
CGCTGGTAGC GGTGGTTTTT GCGACCATCG CCACCAAAAA				7020 7020



TCAAGAAGAT CCTTTGA					7080
AGTTCTTCTA GGAAACTA	AGA AAAGATGCCC	CAGACTGCGA	GTCACCTTGC	TTTTGAGTGC	7080
TTAAGGGATT TTGGTCA	rga gattatcaaa	AAGGATCTTC	ACCTAGATCC	TTTTAAATTA	7140
AATTCCCTAA AACCAGTA	ACT CTAATAGTTT	TTCCTAGAAG	TGGATCTAGG	AAAATTTAAT	7140
AAAATGAAGT TTGCGGC					7200
TTTTACTTCA AACGCCGG					7200
GTTACCAATG CTTAATCA					7260
CAATGGTTAC GAATTAG			1		7260
TAGTTGCCTG ACTCCCCC					7320
ATCAACGGAC TGAGGGG		·			7320
CCAGTGCTGC AATGATAC					7380
GGTCACGACG TTACTATO			•		7380
ACCAGCCAGC CGGAAGGC					7440
TGGTCGGTCG GCCTTCCC					7440
AGTCTATTAA TTGTTGCC					7500
TCAGATAATT AACAACGG				,	7500
ACGTTGTTGC CATTGCTA					7560
TGCAACAACG GTAACGAT					7560
TCAGCTCCGG TTCCCAAC					7620
AGTCGAGGCC AAGGGTTC					7620
CGGTTAGCTC CTTCGGTC					7680
GCCAATCGAG GAAGCCAG					7680
TCATGGTTAT GGCAGCAC					7740
AGTACCAATA CCGTCGTC	!		•		7740
CTGTGACTGG TGAGTACT		= "			7800
GACACTGACC ACTCATGA	AGT TGGTTCAGTA	AGACTCTTAT	CACATACGCC	GCTGGCTCAA	7800



pICAST OMC

GCTCTTGCCC GGCGTCAA	TA CGGGATAATA	CCGCGCCACA	TAGCAGAACT	TTAAAAGTGC	7860
CGAGAACGGG CCGCAGTT	AT GCCCTATTAT	GGCGCGGTGT	ATCGTCTTGA	AATTTTCACG	7860
TCATCATTGG AAAACGTT	CT TCGGGGCGAA	AACTCTCAAG	GATCTTACCG	CTGTTGAGAT	7920
AGTAGTAACC TTTTGCAA	GA AGCCCCGCTT	TTGAGAGTTC	CTAGAATGGC	GACAACTCTA	7920
CCAGTTCGAT GTAACCCA	CT CGTGCACCCA	ACTGATCTTC	AGCATCTTTT	ACTTTCACCA	7980
GGTCAAGCTA CATTGGGT	GA GCACGTGGGT	TGACTAGAAG	TCGTAGAAAA	TGAAAGTGGT	7980
GCGTTTCTGG GTGAGCAA	AA ACAGGAAGGC	AAAATGCCGC	AAAAAAGGGA	ATAAGGGCGA	8040
CGCAAAGACC CACTCGTT	•				8040
CACGGAAATG TTGAATAC	TC ATACTCTTCC	TTTTTCAATA	TTATTGAAGC	ATTTATCAGG	8100
GTGCCTTTAC AACTTATG					8100
GTTATTGTCT CATGAGCG			,		8160
CAATAACAGA GTACTCGC	CT ATGTATAAAC	TTACATAAAT	CTTTTTATTT	GTTTATCCCC	8160
TTCCGCGCAC ATTTC	·				8175
AAGGCGCGTG TAAAG					8175

FIG.12L



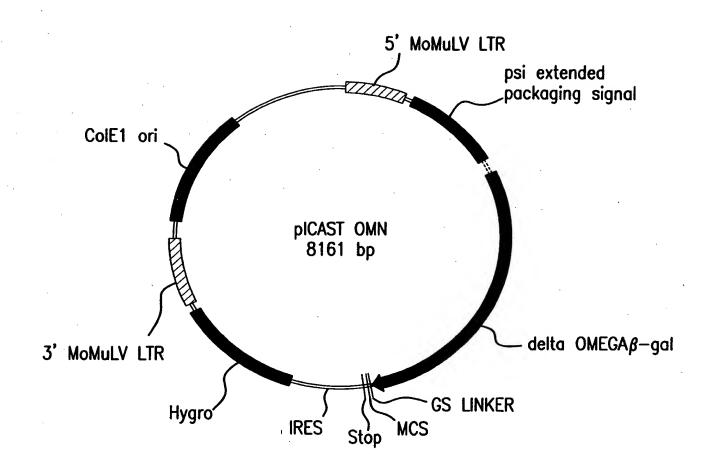


FIG.13A



	AATATGGGCC TTATACCCGG					60 60
	CAGATGGAAC GTCTACCTTG					120 120
	GCTCAGGGCC CGAGTCCCGG					180 180
	ACCATCAGAT TGGTAGTCTA					240 240
	TAACCAATCA ATTGGTTAGT	•				300 300
	AGAGCCCACA TCTCGGGTGT	•				360 360
	CCGTGTATCC GGCACATAGG	· · · · · · · · · · · · · · · · · · ·				420 420
ACAAGGAACC	GAGGGTCTCC CTCCCAGAGG	AGACTGACTA	ACTGATGGGC	AGTCGCCCCC	AGAAAGTAAA	480 480
CCCCGAGCA	CCGGGATCGG GGCCCTAGCC	CTCTGGGGAC	GGGTCCCTGG	TGGCTGGGTG	GTGGCCCTCC	540 540
GTTCGACCGG	AGCAACTTAT TCGTTGAATA	GACACAGACA	GGCTAACAGA	TCACAGATAC	TGACTAAAAT	60 0 60 0
ACGCGGACGC	TCGGTACTAG AGCCATGATC	AATCGATTGA	TCGAGACATA	GACCGCCTGG	GCACCACCTT	660 660
GACTGCTCAA	CTGAACACCC GACTTGTGGG	CCGGCGTTGG	GACCCTCTGC	AGGGTCCCTG	AAACCCCCGG	720 720
	CCCGACCTGA GGGCTGGACT					780 780



	AGGAGACGAG TCCTCTGCTC					840 840
	CCGAAGCCGC GGCTTCGGCG					900 900
	TTCTGTATTT AAGACATAAA					960 960
	GTAACTGGAA CATTGACCTT					1020 1020
	GTTGGGTTAC CAACCCAATG					1080 1080
	GCACCTTTAA CGTGGAAATT					1140` 1140
	ATGGACACCC TACCTGTGGG					1200 1200
AAACTGGGGG	CTCCCTGGGT GAGGGACCCA	GTTCGGGAAA	CATGTGGGAT	TCGGAGGCGG	AGGAGAAGGA	1260 1260
GGTAGGCGGG	CGTCTCTCCC GCAGAGAGGG	GGAACTTGGA	GGAGCAAGCT	GGGGCGAGC	TAGGAGGAA	1320 1320
ATAGGTCGGG	TCACTCCTTC AGTGAGGAAG	AGATCCGCGG	CCGGCGAGAT	CGGGTAATTA	TGCTGAGTGA	1380 1380
TATCCCGCTA	AGCTTGTGGT	ACGTGGTAGT	AGTAGTAGTG	CAGCTGCTTG		1440 1440
AAGGCTTCTT	GACCTACTCG CTGGATGAGC	TCTACCCGCA	CTAATGCCTA	AGTGACCGGC	AGCAAAATGT	1500 1500
	TGGGAAAACC ACCCTTTTGG					1560 1560



	TGGCGTAATA ACCGCATTAT					1620 1620
	GGCGAATGGC CCGCTTACCG					1680 1680
	TGCGATCTTC ACGCTAGAAG					1740 1740
	GATGCGCCCA CTACGCGGGT					1800 1800
	ACGGAGAATC TGCCTCTTAG					1860 1860
	GAAGGCCAGA CTTCCGGTCT					1920 1920
	GGGCGCTGGG CCCGCGACCC					1980 1980
	TTTTTACGCG AAAAATGCGC					2040 2040
	TATCTGGAAG ATAGACCTTC					2100 2100
	CATAAACCGA GTATTTGGCT		•			2160 2160
	AGCCGCGCTG TCGGCGCGAC				AGTTGCGTGA TCAACGCACT	2220 2220
GATGGATGCC	GTAACAGTTT CATTGTCAAA	GAAATACCGT	CCCACTTTGC	GTCCAGCGGT	CGCCGTGGCG	2280 2280
	GGTGAAATTA CCACTTTAAT					2340 2340



	GAAAACCCGA CTTTTGGGCT					2400 2400
	CACACCGCCG GTGTGGCGGC					2460 2460
	CGGATTGAAA GCCTAACTTT					2520 2520
	CGTCACGAGC GCAGTGCTCG					2580 2580
	ATCCTGCTGA TAGGACGACT					2640 2640
	CCGCTGTGGT GGCGACACCA					2700 2700
	GAAACCCACG CTTTGGGTGC					2760 2760
CGATGGCCGC	ATGAGCGAAC TACTCGCTTG	CGCATTGCGC	TTACCACGTC	GCGCTAGCAT	TAGTGGGCTC	2820 2820
ACACTAGTAG	TGGTCGCTGG ACCAGCGACC	CCTTACTTAG	TCCGGTGCCG	CGATTAGTGC	TGCGCGACAT	2880 2880
AGCGACCTAG	AAATCTGTCG TTTAGACAGC	TAGGAAGGGC	GGGCCACGTC	ATACTTCCGC	CGCCTCGGCT	2940 2940
GTGGTGCCGG	ACCGATATTA TGGCTATAAT	AAACGGGCTA	CATGCGCGCG	CACCTACTTC	TGGTCGGGAA	3000 3000
GGGCCGACAC	CCGAAATGGT GGCTTTACCA	GGTAGTTTT	TACCGAAAGC	GATGGACCTC	TCTGCGCGGG	3060 3060
	TGCGAATACG ACGCTTATGC				,	3120 3120



				TTCGTCTGGG AAGCAGACCC	3180 3180
			and the second s	TCGGCTTACG AGCCGAATGC	3240 3240
	· ·			CTGGTCTTTG GACCAGAAAC	 3300 3300
				TTTTTCCAGT AAAAAGGTCA	 3360 3360
				CATAGCGATA GTATCGCTAT	3420 3420
				GGTGAAGTGC CCACTTCACG	 3480 3480
				CCGCAGCCGG GGCGTCGGCC	 3540 3540
				ACCGCATGGT TGGCGTACCA	 3600 3600
		·		CTCAGTGTGA GAGTCACACT	 3660 3660
				GATTTTTGCA CTAAAAACGT	 3720 3720
•				TCACAGATGT AGTGTCTACA	 3780 3780
				CGTGTCGATA GCACAGCTAT	3840 3840
				AATTGACCGG TTAACTGGCC	 3900 3900



TAGATAAGTG	ACTGATTAGA	TGCATTTCGA	CTAGATCCCT	CGACCAATTC	CGGTTATTTT	3960
ATCTATTCAC	TGACTAATCT	ACGTAAAGCT	GATCTAGGGA	GCTGGTTAAG	GCCAATAAAA	3960
•		GGCAATGTGA				4020
		CCGTTACACT				4020
		TCCCCTCTCG				4080
		AGGGGAGAGC				4080
		GAAGCTTCTT				4140
		CTTCGAAGAA				4140
		CCTGGCGACA				4200
		GGACCGCTGT				4200
		GCACAACCCC				4260
		CGTGTTGGGG				4260
		TCAAGCGTAT				4320
		AGTTCGCATA				4320
		GATCTGGGGC			•	4380
		CTAGACCCCG				4380
		GCCCCCGAA				4440
		CGGGGGGCTT				4440
		AAGCCTGAAC				4500
		TTCGGACTTG			•	4500
		TCCGACCTGA				4560
		AGGCTGGACT				4560
		GGGCGTGGAT				4620
		CCCGCACCTA				4620
		GTTTATCGGC				4680
CAAAGATGTT	ICTAGCAATA	CAAATAGCCG	IGAAACGTAG	CCGGCGCGAG	GGCTAAGGCC	4680



CATTGGGGAA GTAACCCCTT		 	-	4740 4740
GTTGCAAGAC CAACGTTCTG		 		4800 4800
GGATGCGATC CCTACGCTAG				4860 4860
AGGAATCGGT TCCTTAGCCA		 		4920 4920
 GTATCACTGG CATAGTGACC		 		4980 4980
TGAGCTGATG ACTCGACTAC				5040 5040
CGGCTCCAAC GCCGAGGTTG		 		5100 5100
GGCGATGTTC CCGCTACAAG				5160 5160
GGCTTGTATG CCGAACATAC	•			5220 5220
GCCGCGGCTC CGGCGCCGAG			-	5280 5280
TGACGGCAAT ACTGCCGTTA		 	TGCGACGCAA ACGCTGCGTT	5340 5340
CGGAGCCGGG GCCTCGGCCC				5400 5400
TGGCTGTGTA ACCGACACAT		 	CCCAGCACTC GGGTCGTGAG	5460 5460



	AAAGGAATAG TTTCCTTATC					5520 5520
	TCCAGAAAAA AGGTCTTTTT					5580 5580
	CGCCATTTTG GCGGTAAAAC					5640 5640
	CAGGAACAGA GTCCTTGTCT					5700 5700
	GCCCCGGCTC CGGGGCCGAG					5760 5760
	TGGTAAGCAG ACCATTCGTC					5820 5820
	GCCCTCAGCA CGGGAGTCGT					5880 5880
	GACCCTGTGC CTGGGACACG					59 40 59 40
	CTGCTCCCCG GACGAGGGGC					6000 6000
	TTGACTGAGT AACTGACTCA					6060 6060
	TGTGGTCTCG ACACCAGAGC					6120 6120
GTCAGCGGGG CAGTCGCCCC	GTCTTTCATT CAGAAAGTAA	CATGCAGCAT GTACGTCGTA	GTATCAAAAT CATAGTTTTA	TAATTTGGTT ATTAAACCAA	TTTTTTCTTA AAAAAAGAAT	6180 6180
	TTAAATGGCC AATTTACCGG					6240 6240



GGTTTGCGTA	TTGGCGCTCT	TCCGCTTCCT	CGCTCACTGA	CTCGCTGCGC	TCGGTCGTTC	6300
	AACCGCGAGA					6300
GGCTGCGGCG	AGCGGTATCA	GCTCACTCAA	AGGCGGTAAT	ACGGTTATCC	ACAGAATCAG	6360
	TCGCCATAGT					6360
GGGATAACGC	AGGAAAGAAC	ATGTGAGCAA	AAGGCCAGCA	AAAGGCCAGG	AACCGTAAAA	6420
	TCCTTTCTTG					6420
	GCTGGCGTTT					6480
	CGACCGCAAA			·		6480
	TCAGAGGTGG					6540
	AGTCTCCACC				•	6540
	CCTCGTGCGC					6600
	GGAGCACGCG					6600
	TTCGGGAAGC					6660
	AAGCCCTTCG					6660
	CGTTCGCTCC					6720
	GCAAGCGAGG					6720
	ATCCGGTAAC		·			6780
	TAGGCCATTG					6780
	AGCCACTGGT					6840
	TCGGTGACCA				•	6840
	GTGGTGGCCT					6900
	CACCACCGGA					6900
	GCCAGTTACC					6960
	CGGTCAATGG					6960
	TAGCGGTGGT					7020
GG I GGCGACC	ATCGCCACCA	AAAAAACAAA	CGTTCGTCGT	CTAATGCGCG	ICTITITIC	7020



	TA CGGGGTCTGA CGCTCAGTGG AACGAAAACT 708 AT GCCCCAGACT GCGAGTCACC TTGCTTTTGA 708
	AT CAAAAAGGAT CTTCACCTAG ATCCTTTTGC 714 TA GTTTTTCCTA GAAGTGGATC TAGGAAAACG 714
	AG TAAACTTGGT CTGACAGTTA CCAATGCTTA 720 TC ATTTGAACCA GACTGTCAAT GGTTACGAAT 720
	GT CTATTTCGTT CATCCATAGT TGCCTGACTC 726 CA GATAAAGCAA GTAGGTATCA ACGGACTGAG 726
	AG GGCTTACCAT CTGGCCCCAG TGCTGCAATG 732 TC CCGAATGGTA GACCGGGGTC ACGACGTTAC 732
	CA GATTTATCAG CAATAAACCA GCCAGCCGGA 738 GT CTAAATAGTC GTTATTTGGT CGGTCGGCCT 738
•	CT TTATCCGCCT CCATCCAGTC TATTAATTGT 7440 GA AATAGGCGGA GGTAGGTCAG ATAATTAACA 7440
	CA GTTAATAGTT TGCGCAACGT TGTTGCCATT 7500 GT CAATTATCAA ACGCGTTGCA ACAACGGTAA 7500
	CG TTTGGTATGG CTTCATTCAG CTCCGGTTCC 7560 GC AAACCATACC GAAGTAAGTC GAGGCCAAGG 7560
	CC ATGTTGTGCA AAAAAGCGGT TAGCTCCTTC 7620 GG TACAACACGT TTTTTCGCCA ATCGAGGAAG 7620
	FG GCCGCAGTGT TATCACTCAT GGTTATGGCA 7680 AC CGGCGTCACA ATAGTGAGTA CCAATACCGT 7680
	CA TCCGTAAGAT GCTTTTCTGT GACTGGTGAG 7740 GT AGGCATTCTA CGAAAAGACA CTGACCACTC 7740
	GT ATGCGGCGAC CGAGTTGCTC TTGCCCGGCG 7800 CA TACGCCGCTG GCTCAACGAG AACGGGCCGC 7800



pICAST OMN

TCAATACGGG ATAATACCGC GCCACATAGC AGAACTTTAA AAGTGCTCAT	CATTGGAAAA	7860
AGTTATGCCC TATTATGGCG CGGTGTATCG TCTTGAAATT TTCACGAGTA	GTAACCTTTT	7860
CGTTCTTCGG GGCGAAAACT CTCAAGGATC TTACCGCTGT TGAGATCCAG	TTCGATGTAA	7920
GCAAGAAGCC CCGCTTTTGA GAGTTCCTAG AATGGCGACA ACTCTAGGTC	AAGCTACATT	7920
CCCACTCGTG CACCCAACTG ATCTTCAGCA TCTTTTACTT TCACCAGCGT	TTCTGGGTGA	7980
GGGTGAGCAC GTGGGTTGAC TAGAAGTCGT AGAAAATGAA AGTGGTCGCA		7980
GCAAAAACAG GAAGGCAAAA TGCCGCAAAA AAGGGAATAA GGGCGACACG		8040
CGTTTTTGTC CTTCCGTTTT ACGGCGTTTT TTCCCTTATT CCCGCTGTGC		8040
ATACTCATAC TCTTCCTTTT TCAATATTAT TGAAGCATTT ATCAGGGTTA		8100
TATGAGTATG AGAAGGAAAA AGTTATAATA ACTTCGTAAA TAGTCCCAAT		8100
AGCGGATACA TATTTGAATG TATTTAGAAA AATAAACAAA TAGGGGTTCC		8160
TCGCCTATGT ATAAACTTAC ATAAATCTTT TTATTTGTTT ATCCCCAAGG	CGCGTGTAAA	8160
C		8161
G	,	8161

FIG.13L



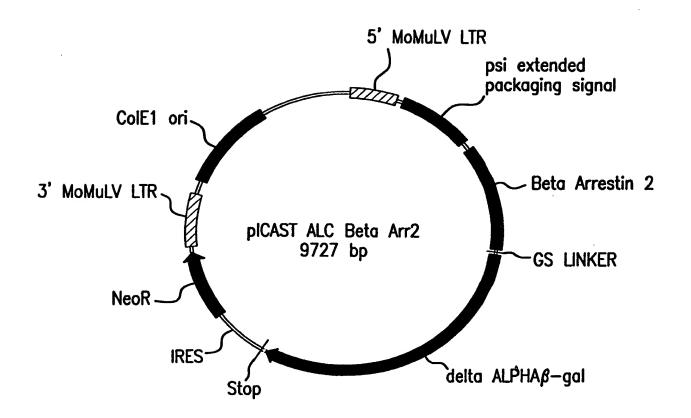


FIG.14



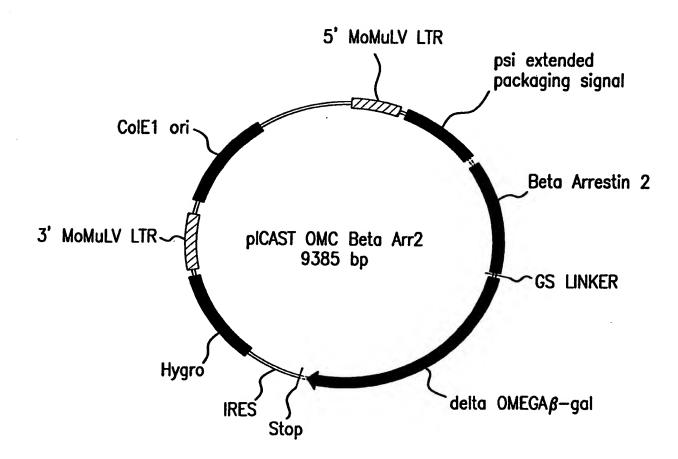


FIG.15



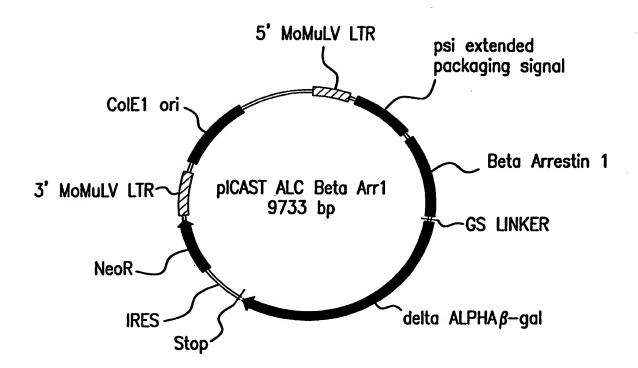


FIG.16



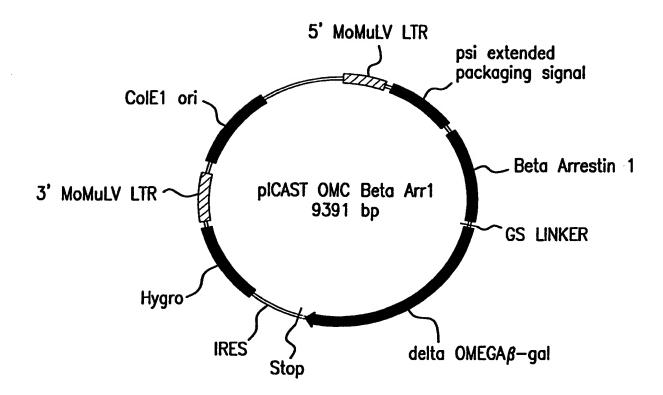


FIG.17



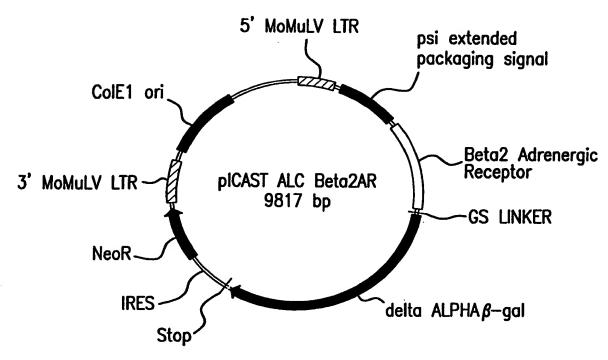


FIG.18



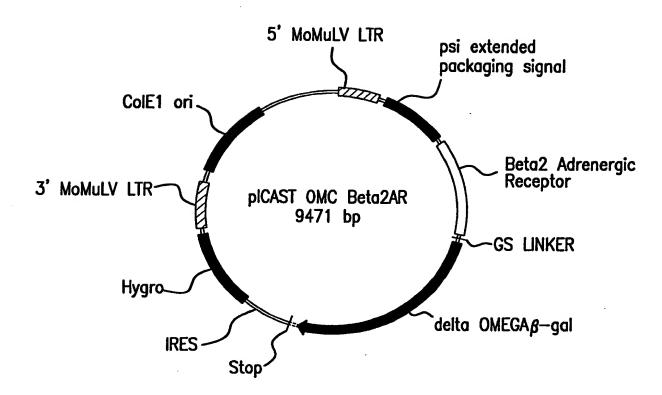


FIG.19



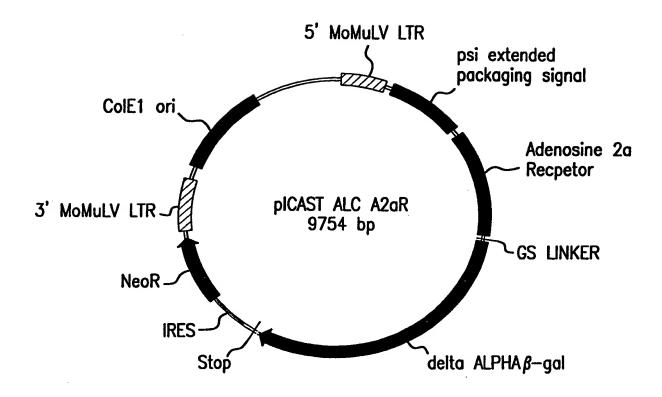


FIG.20



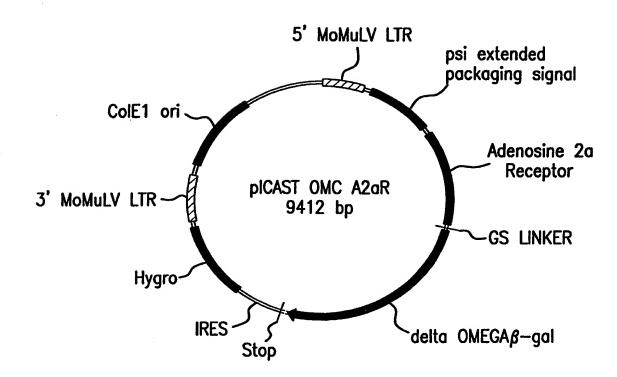


FIG.21



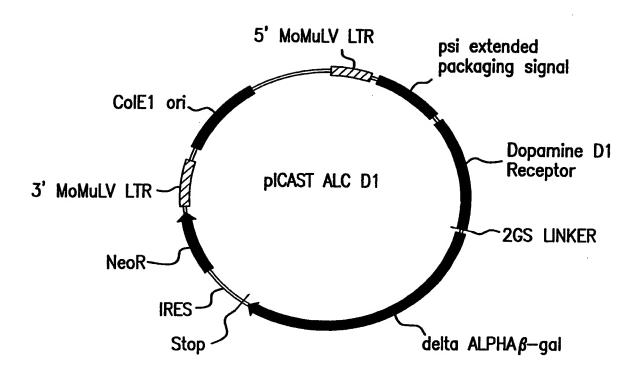
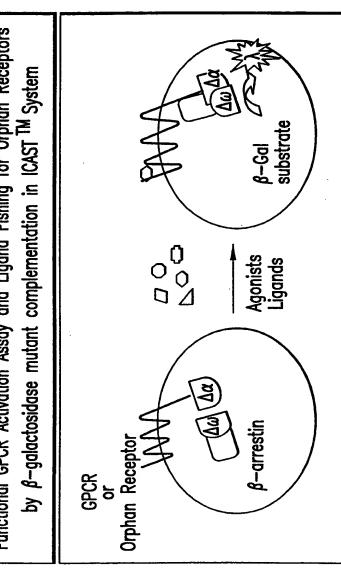


FIG.22



Functional GPCR Activation Assay and Ligand Fishing for Orphan Receptors by β -galactosidase mutant complementation in ICAST TM System



Examples

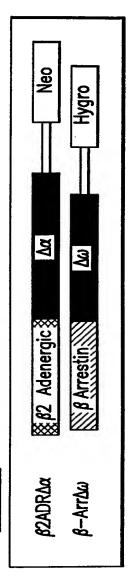


FIG. 23